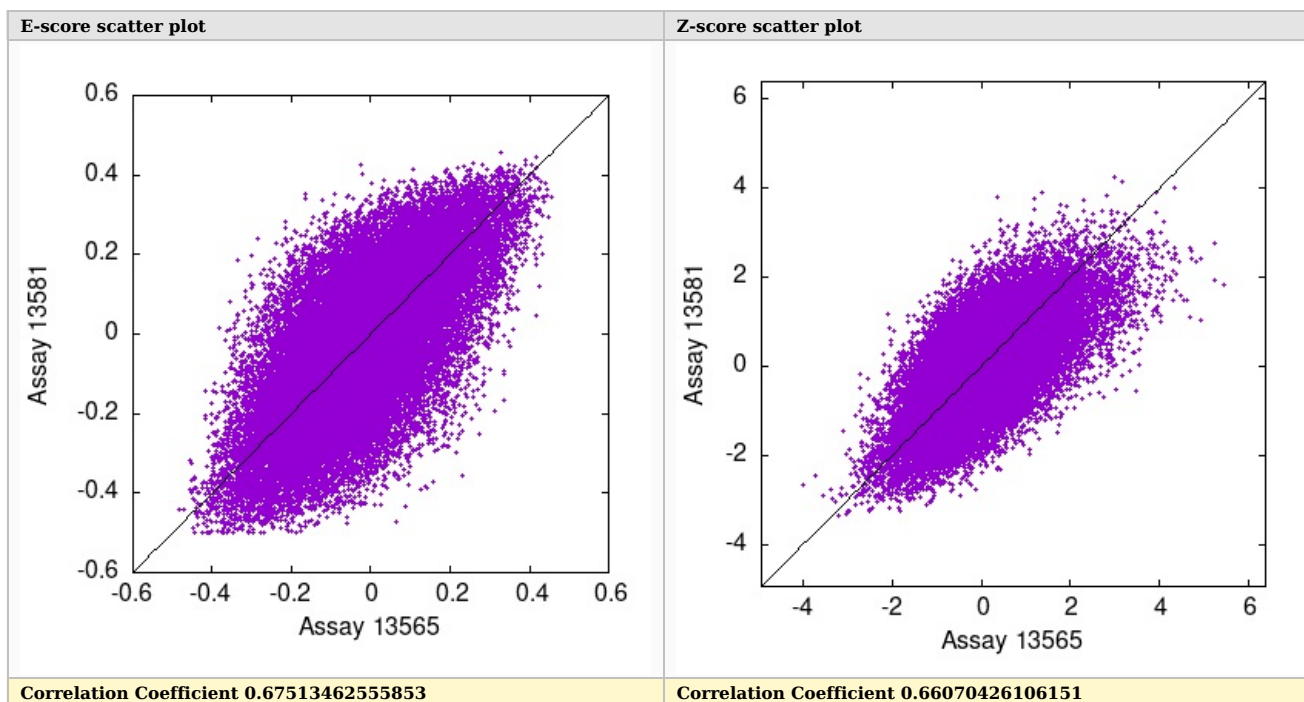


QC report



Top scoring motifs for Assay 13565

Protein ID: pTH13957.1 Gene: CHAMP1.DBD Domain: zf-C2H2 Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment
ATAATATA	0.45767	---TATATTAT---
ACTAATAA	0.44891	TTATTAGT----
AACTAACG	0.44567	--CGTTAGTT---
GTTAGTTA	0.44439	--GTTAGTTA--
AACTAACT	0.44214	-AGTTAGTT---
AAAGTAAT	0.43872	--ATTACTTT--
GTTAACCTA	0.43644	----TAGTTAAC
ATTAGTTG	0.43482	--ATTAGTTG--
AAGTAACA	0.43339	-TGTTACTT---
AACTAATG	0.43212	-CATTAGTT---

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment
ACTAATAA	5.45107	TTATTAGT----
GTTAGTTA	5.26544	--GTTAGTTA--
ATAATATA	5.25693	---TATATTAT---
AACTAACG	4.94130	-CGTTAGTT---
ACTAACGG	4.92239	CCGTTAGT----
ATATGTTA	4.85238	--ATATGTTA--
AAGTAATT	4.74237	-AATTACTT---
GTTAACTA	4.73338	----TAGTTAAC
AACTAACT	4.70248	-AGTTAGTT---
CGTTAGTC	4.68527	-CGTTAGTC---

Top scoring motifs for Assay 13581

Protein ID: pTH13957.2 Gene: CHAMP1.DBD Domain: zf-C2H2 Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment
AATAATAA	0.45602	--AATAATAA--
ACTAATTA	0.44618	---ACTAATTA---
ACTATAAT	0.43729	---ACTATAAT---
ATATTAAG	0.42970	---ATATTAAG---
ACTAACAT	0.42635	--ACTAACAT---
TAATAATA	0.42584	-TAATAATA---
ATATAACA	0.42535	-ATATAACA---
ATTATTTT	0.42493	GAAATAAT---
AGTAATTA	0.42441	--AGTAATTA---
AAATAATA	0.42221	-AAATAATA---

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment
AATAATAA	4.25128	--AATAATAA--
AGTAATTA	4.13176	--AGTAATTA---
ACTAATTA	4.00186	---ACTAATTA---
ACTATAAT	3.92271	---ACTATAAT---
ATATCACA	3.91826	---ATATCACA---
ATATAACA	3.82133	-ATATAACA---
ATTATTTT	3.78685	GAAATAAT---
ATATACAA	3.78323	--ATATACAA---
ATATTAAG	3.72001	---ATATTAAG---
ATTAATCA	3.63989	--ATTAATCA---